REMARKS

Applicant requests entry of the instant amendment prior to examination of the above-identified patent application on the merits.

Claims 1-12 and 14-16 are pending in the above-identified application. Claim 13 has been cancelled. Claims 1-12 have been amended. New claims 14-16 have been added by the instant amendment.

Attached hereto is a marked-up version of the changes made by the instant amendment. The attached page is captioned "VERSION WITH MARKINGS TO SHOW CHANGES MADE" and is only included for the Examiner's convenience. Should any discrepancies be discovered, the version presented in the preceding "IN THE CLAIMS" section shall be deemed to be correct.

Applicants assert that these new claims are fully supported by the application as filed and do not constitute new matter.

Applicants submit herewith a Substitute Sequence Listing in paper and computer form. I hereby state that the content of the paper and computer readable copies of the Substitute Sequence Listing submitted in accordance with 37 C.F.R. §1.821(c) and (e), are the same. I hereby state that the content of the paper and computer readable copies of the Substitute Sequence Listing, submitted in accordance with 37 C.F.R. §1.821(g), herein does not include new matter.

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The Commissioner is hereby authorized to charge any fees due with this submission not otherwise enclosed herewith to Deposit Account No. 02-4377. Please credit any overpayment of fees associated with this filing to the above-identified deposit account. A duplicate of this page is enclosed.

Respectfully submitted,

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In the following sections, added text is marked with double underlining. e.g. added text, and deleted text is marked by a single strikethrough, e.g. deleted text.

VERSION WITH MARKINGS TO SHOW CHANGES MADE

IN THE CLAIMS

Claims 1-10 have been amended as follows:

- 1. A method for producing <u>an</u> isotransgenic plant lines, line comprising the following steps of:
 - a) transforming the plant cells of a plant-hybrid consisting of plant, the crossing of two parental lines, line of which are a line of interest and a line suited to transformation, with a vector carrying comprising a T-DNA containing a transgene;
 - b) selecting-the hybrid primary transformants which have integrated said T-DNA only, into the genome of the line of interest;
 - c) backcrossing, selected hybrid primary transformants with the parental line of interest, said primary transformants selected in b), and
 - d) ____selecting the individuals at least one transgenic individual derived from these each backcrosses backcross until the isotransgenic lines line are is produced.
- 2. The method as claimed inof Claim 1, characterized in that wherein the step for selecting the hybrid primary transformants consists incomprises identifying the genomic sequences adjacent to the T-DNA inserted, in order to determine and determining the parent genome which has received said T-DNA.
- 3. The method as claimed inof Claim 2, inwherein which determining the determination of the parent plant genome which has received said T-DNA, using said genomic

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sequences adjacent to the T-DNA; is carried out according to a technique selected from the group consisting of an RFLP technique or and a sequencing method.

- 4. The method as claimed in one of <u>Claim 1</u>, wherein the <u>Claims 1 to 3</u>, individual selected in which the individuals in which the (d) has
 - a chromosome which has received having the T-DNA has conserved but otherwise having a genotype geneotype entirely of the line of interest type; and which have
 - b) are selected from the first backcross in c).
- 5. The method as claimed in one of the preceding claims, characterized in that it ecomprises a subsequent step of Claim 1 further comprising crossing between the isotransgenic plant line according to the invention and another a second line of interest, in particular another isotransgenic line containing a different transgene, for producing a hybrid line.
- 6. The method as claimed in one of <u>Claim 1, wherein</u> the <u>preceding claims</u>, characterized in that the <u>hybrid</u> plant cells is originate selected from the group consisting of a large crop species chosen from maize <u>plant</u>, wheat, rapeseed, sunflower, pea, soybean and barley, or from a vegetable or <u>plant</u>, and floral species <u>plant</u>.
- 7. The method as claimed in one of the preceding claims, characterized in that <u>Claim</u> 1, wherein the T-DNA comprises in particular a nucleotide sequence encoding a protein which confers agronomic properties and/or properties of resistance to diseases.
- 8. The method as claimed in one of the preceding claims, characterized in that Claim 1, wherein the isotransgenic lines produced are line is a commercial elite lines line.
- 9. The use of the The method as claimed in one of Claims 1 to 8, characterized in that it allows Claim 1, wherein the introgression of several transgenic characteristics into a isogenic plant, without adding is substantially free of fragments linked to the transgene which may be the subject of a genetic burden.

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- 10. A method which makes it possible to target of identifying the parent genome which has received a T-DNA after transformation of a hybrid, comprising the identification of the identifying genomic sequences adjacent to the T-DNA inserted.
- 11. A transgenic plant, organ or part of a plant, in particular seed, obtained according to by the invention in one or other method of the steps described in Claim 1 or 5.
- 12. A true<u>An</u> isotransgenic line produced from hybrid transformants as claimed in one by the method of Claims 1 or 7, characterized in that they have fixed wherein the isotransgenic line has a pure line of interest genotype over the entire genome and have stably integrated the T-DNA containing the transgene.